

OIPE

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/993,179

DATE: 12/05/2001  
TIME: 09:54:30

Input Set : A:\sequence listing.txt  
Output Set: N:\CRF3\11212001\I993179.raw

Does Not Comply  
Corrected Diskette Needed

gr 1-3

3 <110> APPLICANT: McCarthy, Sean A.  
4 Kuranda, Michael Joseph  
5 Bulawa, Christine Ellen  
6 Bossone, Steven  
8 <120> TITLE OF INVENTION: METHOD FOR IDENTIFYING GENES ENCODING SIGNAL SEQUENCES  
10 <130> FILE REFERENCE: 09404/032001  
12 <140> CURRENT APPLICATION NUMBER: US/09/993,179  
13 <141> CURRENT FILING DATE: 2001-11-06  
15 <160> NUMBER OF SEQ ID NOS: 15  
17 <170> SOFTWARE: FastSEQ for Windows Version 3.0

## ERRORRED SEQUENCES

52 <210> SEQ ID NO: 2  
53 <211> LENGTH: 50  
54 <212> TYPE: PRT  
55 <213> ORGANISM: Homo sapiens  
57 <400> SEQUENCE: 2  
58 Met Lys Gly Thr Cys Val Ile Ala Trp Leu Phe Ser Ser Leu Gly Leu  
E--> 59 1 5 5 10 15 10 15  
60 Trp Arg Leu Ala His Pro Glu Ala Gln Gly Thr Thr Gln Cys Gln Arg  
E--> 61 20 25 30  
62 Thr Leu Glu Val Asn Ile Val Ser Pro Ser Ser Lys Ala Thr Phe Ser  
E--> 63 35 40 45  
64 Pro Ser  
65 50  
112 <210> SEQ ID NO: 4  
113 <211> LENGTH: 125  
114 <212> TYPE: PRT  
115 <213> ORGANISM: Homo sapiens  
117 <400> SEQUENCE: 4  
118 Met Arg Ser Leu Leu Arg Thr Pro Phe Leu Cys Gly Leu Leu Trp Ala  
E--> 119 1 5 10 15  
120 Phe Cys Ala Pro Gly Ala Arg Ala Glu Glu Pro Ala Ala Ser Phe Ser  
E--> 121 20 25 30  
122 Gln Pro Gly Ser Met Gly Leu Asp Lys Asn Thr Val His Asp Gln Glu  
E--> 123 35 40 45  
124 His Ile Met Glu His Leu Glu Gly Val Ile Asn Lys Pro Glu Ala Glu  
E--> 125 50 55 60  
126 Met Ser Pro Gln Glu Leu Gln Leu His Tyr Phe Lys Met His Asp Tyr  
E--> 127 65 70 75 80  
128 Asp Gly Asn Asn Leu Leu Asp Gly Leu Glu Leu Ser Thr Ala Ile Thr  
E--> 129 85 90 95  
130 His Val His Lys Glu Gly Ser Glu Gln Ala Pro Leu Glu Val Asn  
E--> 131 100 105 110  
132 Ile Val Ser Pro Ser Ser Lys Ala Thr Phe Ser Pro Ser

misaligned  
and id nos  
(see item 3  
in Error Summary  
sheet)

same  
err

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/993,179

DATE: 12/05/2001  
TIME: 09:54:30

Input Set : A:\sequence listing.txt  
Output Set: N:\CRF3\11212001\I993179.raw

E--> 133 115 120 125  
135 <210> SEQ ID NO: 5  
136 <211> LENGTH: 32  
137 <212> TYPE: PRT  
138 <213> ORGANISM: Mus musculus  
140 <400> SEQUENCE: 5  
141 Met Lys Gly Ala Cys Ile Leu Ala Trp Leu Phe Ser Ser Leu Gly Val  
E--> 142 1 5 10 15  
143 Trp Arg Leu Ala Arg Pro Glu Thr Gln Asp Pro Ala Lys Cys Gln Arg  
E--> 144 20 25 30  
146 <210> SEQ ID NO: 6  
147 <211> LENGTH: 45  
148 <212> TYPE: PRT  
149 <213> ORGANISM: Homo sapiens  
151 <400> SEQUENCE: 6  
152 Met Ser Pro Gln Glu Leu Gln Leu His Tyr Phe Lys Met His Asp Tyr  
E--> 153 1 5 10 15  
154 Asp Gly Asn Asn Leu Leu Asp Gly Leu Glu Leu Ser Thr Ala Ile Thr  
E--> 155 20 25 30  
156 His Val His Lys Glu Glu Gly Ser Glu Gln Ala Pro Leu  
E--> 157 35 40 45  
238 <210> SEQ ID NO: 14  
239 <211> LENGTH: 32  
240 <212> TYPE: PRT  
241 <213> ORGANISM: Homo sapiens  
243 <400> SEQUENCE: 14  
244 Met Lys Gly Thr Cys Val Ile Ala Trp Leu Phe Ser Ser Leu Gly Leu  
E--> 245 1 5 10 15  
246 Trp Arg Leu Ala His Pro Glu Ala Gln Gly Thr Thr Gln Cys Gln Arg  
E--> 247 20 25 30  
249 <210> SEQ ID NO: 15  
250 <211> LENGTH: 108  
251 <212> TYPE: PRT  
252 <213> ORGANISM: Homo sapiens  
254 <400> SEQUENCE: 15  
255 Met Arg Ser Leu Leu Arg Thr Pro Phe Leu Cys Gly Leu Leu Trp Ala  
E--> 256 1 5 10 15  
257 Phe Cys Ala Pro Gly Ala Arg Ala Glu Glu Pro Ala Ala Ser Phe Ser  
E--> 258 20 25 30  
259 Gln Pro Gly Ser Met Gly Leu Asp Lys Asn Thr Val His Asp Gln Glu  
E--> 260 35 40 45  
261 His Ile Met Glu His Leu Glu Gly Val Ile Asn Lys Glu Ala Glu Met  
E--> 262 50 55 60  
263 Ser Pro Gln Glu Leu Gln Leu His Tyr Phe Lys Met His Asp Tyr Asp  
E--> 264 65 70 75 80  
265 Gly Asn Asn Leu Leu Asp Gly Leu Glu Leu Ser Thr Ala Ile Thr His  
E--> 266 85 90 95  
267 Val His Lys Glu Glu Gly Ser Glu Gln Ala Pro Leu  
E--> 268 100 105

same

same

same

same

(see next page)

09/993,179

3

<400> 1			
ggggaccgtg tttgtggccc ccaagccgtt gccccccatt ttggaactca gcgagtaggg			60
ggcggtctcg gggaaagtggc agggggcgca gcagctgctg cctccacttc cctagccagg			120
tgcgtaaagag gatttcgga gcccgtctgg ccccccaggcg ctggatgact ggcaccagcg			180
ctccctcgac ac tctgtttgtt gtgtgagact tgggtggag tgccacgtg gctgtggagt			240
cagtgattt catgatttag gaaacgcgtc ctccatcctc tcttccttg gcactttcca			300
catatgagga gaagaagagc ttctgttttag aagacacgtg cccagagtca gaggccccctt			360
gcccacc atg aag gga acc tgg ttttata gca tgg ctg ttc tca agc ctg	409		
Met Lys Gly Thr Cys Val Ile Ala Trp Leu Phe Ser Ser Leu →			
1 5 10			
ggg ctg tgg aga ctc gcc cac cca gag gcc cag ggt acg act cag tgc	457		
Gly Leu Trp Arg Leu Ala His Pro Glu Ala Gln Gly Thr Thr Gln Cys			
15 20 25 30			
cag aga aca ctc gag gtg aat att gtt tcc ccc agc tcc aag gca aca	505		
Gln Arg Thr Leu Glu Val Asn Ile Val Ser Pro Ser Ser Lys Ala Thr			
35 40 45			
ttc agt cca agt	517		
Phe Ser Pro Ser			
50			

move amino acid directly  
under /  
respective  
amino acid

(please correct this misalignment)  
in subsequent coding sequences

*fyi*  
Use of n and/or Xaa has been detected in the Sequence Listing.  
Review the Sequence Listing to insure a corresponding  
explanation is presented in the <220> to <223> fields of  
each sequence using n or Xaa.

VERIFICATION SUMMARY  
PATENT APPLICATION: US/09/993,179

DATE: 12/05/2001  
TIME: 09:54:31

Input Set : A:\sequence listing.txt  
Output Set: N:\CRF3\11212001\I993179.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application Number  
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:41 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:45 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:49 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:59 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:2  
M:332 Repeated in SeqNo=2  
L:81 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:3  
L:85 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:3  
L:89 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:3  
L:93 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:3  
L:97 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:3  
L:101 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:3  
L:105 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:3  
L:109 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:3  
L:119 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:4  
M:332 Repeated in SeqNo=4  
L:142 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:5  
M:332 Repeated in SeqNo=5  
L:153 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6  
M:332 Repeated in SeqNo=6  
L:209 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:11  
L:214 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11  
L:245 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:14  
M:332 Repeated in SeqNo=14  
L:256 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:15  
M:332 Repeated in SeqNo=15